Changed a lile from non-ASCII to ASCII Changed the margins in cases where the sequence text was "vrapped" down to the no Edited a format error in the Current Application Data section, specifically: Edited the Current Application Data section with the actual current number. The number applicant was the prior application data; or other Added the mandatory heading and subheadings for "Current Application Data". Edited the "Number of Sequences" field. The applicant spelled out a number instead of Changed the spelling of a mandatory field (the headings or subheadings), specifically: Corrected the SEO ID NO when obviously incorrect. The sequence numbers that were defined or corrected a nucleic number at the end of a nucleic line. SEO ID NO's edited:	xt line. #2
Changed a file from non-ASCII to ASCII Changed the margine in cases where the sequence text was "wrapped down to the ne Edited a format error in the Current Application Data section, specifically: Edited the Current Application Data section with the actual current number. The number applicant was the prior application data; or other formation Data. Edited the "Number of Sequences" field. The applicant spelled out a number instead of Changed the spelling of a mandatory field (the headings or subheadings), specifically: Corrected the SEO ID NO when obviously incorrect. The sequence numbers that were considered the sequence numbers that were considered the second sequences.	xt line. #2
Changed the margins in cases where the sequence text was "wrapped" down to the need to the form of the Current Application Data section. Specifically: Edited the Current Application Data section with the actual current number. The number applicant was the prior application data; or the other the prior application data; or the current Application Data. Added the mandatory heading and subheadings for "Current Application Data". Edited the "Number of Sequences" field. The applicant spelled out a number instead of Changed the spelling of a mandatory field (the headings or subheadings), specifically: Corrected the SEO ID NO when obviously incorrect. The sequence numbers that were defined to the sequence numbers that were defined to the sequence numbers of the seque	xt line. #2
Changed the margins in cases where the sequence text was "wrapped" down to the need to the form of the Current Application Data section. Specifically: Edited the Current Application Data section with the actual current number. The number applicant was the prior application data; or the other the prior application data; or the current Application Data. Added the mandatory heading and subheadings for "Current Application Data". Edited the "Number of Sequences" field. The applicant spelled out a number instead of Changed the spelling of a mandatory field (the headings or subheadings), specifically: Corrected the SEO ID NO when obviously incorrect. The sequence numbers that were defined to the sequence numbers that were defined to the sequence numbers of the seque	xt line. #2
Edited a format error in the Current Application Data section, specifically: Edited the Current Application Data section with the actual current number. The number applicant was the prior application data; or other Added the mandatory heading and subheadings for "Current Application Data". Edited the "Number of Sequences" field. The applicant spelled out a number instead of Changed the spelling of a mandatory field (the headings or subheadings), specifically: Corrected the SEO ID NO when obviously incorrect. The sequence numbers that were defined to the sequence numbers of the sequence numbers o	er inputted by the
Edited the Current Application Data section with the actual current number. The number applicant was the prior application data; or other Added the mandatory heading and subheadings for "Current Application Data". Edited the "Number of Sequences" field. The applicant spelled out a number instead of Changed the spelling of a mandatory field (the headings or subheadings), specifically: Corrected the SEO ID NO when obviously incorrect. The sequence numbers that were detailed.	
Added the mandatory heading and subheadings for "Current Application Data". Edited the "Number of Sequences" field. The applicant spelled out a number instead of Changed the spelling of a mandatory field (the headings or subheadings), specifically: Corrected the SEO ID NO when obviously incorrect. The sequence numbers that were desired.	
Edited the 'Number of Sequences' field. The applicant spelled out a number instead of Changed the spelling of a mandatory field (the headings or subheadings), specifically: Corrected the SEO ID NO when obviously incorrect. The sequence numbers that were desired to the sequence numbers of the	using an inlogor
Changed the spelling of a mandatory field (the headings or subheadings), specifically: Corrected the SEO ID NO when obviously incorrect. The sequence numbers that were experienced in the sequence numbers in the sequence numbers of the sequence numbers in the sequence n	using an intogor
Corrected the SEO ID NO when obviously incorrect. The sequence numbers that were o	
Inserted or corrected a nucleic number at the end of a nucleic line. SEO ID NO's edited:	dited were:
Corrected subheading placement. All responses must be on the same line as each subheapplicant placed a response below the subheading, this was moved to its appropriate place.	eading. Îl Iho co.
Inserted colons after headings/subheadings. Headings edited included: •, . * *	
Deleted extra, invalid, headings-used by an applicant, specifically:	
Deletod: non-ASCII *garbago* at the beginning/end of files: secretary initials/filer page numbers throughout text; other invalid text, such as	name at end of fi
Inserted mandatory headings, specifically:	
Corrected an obvious error in the response, specifically:	
Edited identifiers where upper case is used but lower case is required, or vice versa.	
Corrected an error in the Number of Sequences field, specifically:	
A "Hard Pago Break" code was inserted by the applicant. All occurrences had to be deleted	ed.
Peloted ending stop codon in amino acid sequences and adjusted the "(A)Length:" field ac up to a Patentin bug). Sequences corrected:	cordingly (error
Other:	
3	

Examiner: The above corrections must be communicated to the applicant in the first Office Aciton. DO NOT send a copy of this form.

3

DATE: 12/07/2001

OIPE

```
TIME: 19:16:25
                     PATENT APPLICATION: US/09/960,706
                     Input Set : N:\jumbos\960706.txt
                     Output Set: N:\CRF3\12072001\1960706.raw
      3 <110> APPLICANT: Munger, William E.
      5 <120> TITLE OF INVENTION: Identifying Drugs for and Diagnosis of Benign Prostatic
Hyperplasia Using
              Gene Expression Profiles
      8 <130> FILE REFERENCE: 44921-5029-01US
C--> 10 <140> CURRENT APPLICATION NUMBER: US/09/960,706
C--> 11 <141> CURRENT FILING DATE: 2001-09-24
     13 <150> PRIOR APPLICATION NUMBER: 60/223,323
     14 <151> PRIOR FILING DATE: 2000-08-07
     16 <150> PRIOR APPLICATION NUMBER: 09/873,319
     17 <151> PRIOR FILING DATE: 2001-06-05
     19 <160> NUMBER OF SEQ ID NOS: 1124
     21 <170> SOFTWARE: PatentIn Ver. 2.1
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     24 <211> LENGTH: 333
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     33 ggcctgagcg cctgggaagt ctcttgaggt tgcaggaatc tccagagaaa cataggcgct 120
     34 gcccagccac caccccgaga acactatttg ggctggagtg tgaccgccga ggtgatcctg 180
     35 gcaggagget ggggttgget cetegactee acaaacaetg aggagtgggt ggggacaeca 240
     36 ttgacaccca cccaaacact ggcagagagg gaaggccctt ccagattctg gggcacatgt 300
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     48 <221> NAME/KEY: unsure
     49 <222> LOCATION: (1)..(426)
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     55 atcagcatat cacatggcga gagagaacaa agagctgcca ggctttttaa agcaaccaga 180
     56 totoatgtga acttagagag tgagaatgta aggttagccg agagaaagga ccagagagag 240
     57 agagececaa gtteaggeaa geetttatta acetgtegge tgeeceetta acagteaagg 300
W--> 58 aaagcagccc caagcttaca gaataggggg tttacattng gggcagggag tttgagggag 360
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     60 tgtgaa
     62 <210> SEQ ID NO: 3
     63 <211> LENGTH: 433
     64 <212> TYPE: DNA
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RAW SEQUENCE LISTING

RAW SEQUENCE LISTING DATE: 12/07/2001 PATENT APPLICATION: US/09/960,706 TIME: 19:16:25

Input Set : N:\jumbos\960706.txt

Output Set: N:\CRF3\12072001\I960706.raw

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     72 acaagtgaaa aagaacttga tteettaata aaatgaetta caagtgeaaa aatggaeeee 120
     73 atatgtgctc tgctttactt acacccaact ctaacctgcg agcatattag gaaaaaagaa 180
     74 gcaagtgcat ccaataaaat tatttgtaac ttaatttgta tgaggcggga caaggttgct 240
     75 tttgatccaa gtctggtatt tcttggttaa cagggtgtag attccaggct ttgtggcaac 300
     76 accacattca tgacctccag agactatagc gtgggaagac acctttacag atcaaggggc 360
     77 cccctggagt cacccttaca ggaatccttc tggcctttgg aatcctcctg gcacagacca 420
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     78 tgtctttggt gat
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     90 agtggaacgt gtcaaacaga aatggtgaca atgagttaga actgcagttg tttcaaggta 120
     91 ctacactatt atttaaaaaa aaaactcaca aaaagaaaaa tgttatcact acaagtagga 180
     92 attagaagag agaaateetg geagtetgte tagaggttaa aacattteat geatttgtga 240
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     106 <223> OTHER INFORMATION: n = a or c or g or t
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     111 ggatatattg ctttgataat tataaagcta tttttttcat tttaaaagta gctaagtttt 180
     112 caacatttca aaaacttttt ccagatette tgtataettt tetgtaggea teetagtgaa 240
     113 acatgtacaa ttcaaatgac cacatgctgg agagccaggc gcgtccccat gcaggcgacg 300
W--> 114 tgggcctctt agaagcagee teetgagntg tgtageetee tgcageeata cagteeegte 360
     115 ctggtcgccc gtctcagcct cggtcctgag cagcctgtgg gaatggtggc gacgggacag 420
W--> 116 tccagaacac tcctggacag cccctagccc cagcccctgc ccgcctctcg ggagagcan 479
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     120 <212> TYPE: DNA
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     123 <220> FEATURE:
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124 <223> OTHER INFORMATION: Genbank Accession No. AA009755

RAW SEQUENCE LISTING DATE: 12/07/2001 PATENT APPLICATION: US/09/960,706 TIME: 19:16:25

Input Set : N:\jumbos\960706.txt

Output Set: N:\CRF3\12072001\I960706.raw

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186 <210> SEQ ID NO: 10

RAW SEQUENCE LISTING DATE: 12/07/2001 PATENT APPLICATION: US/09/960,706 TIME: 19:16:25

Input Set : N:\jumbos\960706.txt

Output Set: N:\CRF3\12072001\I960706.raw

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W--> 244 cagectecta aaagtgetgg ggattacagg catgagecae canaceeggg ceaagttett 480

245 tt

RAW SEQUENCE LISTING DATE: 12/07/2001 PATENT APPLICATION: US/09/960,706 TIME: 19:16:25

Input Set : N:\jumbos\960706.txt

Output Set: N:\CRF3\12072001\I960706.raw

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     258 ggagagtaat acaaaaaat aaggaataaa agctaaagat ctaactactc cgaccttcac 180
     259 aattocaget acttgataat aataggagta acccaatgaa tactgtatgg tetgaaaget 240
     260 actatacaat atgattetta acgagaaggg aagggaatta gagaetgtea caaageeetg 300
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     262 ggaagtgatt ctg
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     280 tttctggggc agactttttc cggggccgat ctttgggaac ggacagaaat tcgggtgcgt 180
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     282 ccccg
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     285 <211> LENGTH: 337
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     295 tttaagagtc catcaaagtg tcatatgtgt taggtgtgaa atggcgacac tgggaattac 180
     296 tgttaataag gggtggctgc agcacggtga ttgttatgag aacatcccca ccgccccact 240
     297 tttgtttgaa gactttcgta ctgaactaca tgttgtttac tttcaacaac gtatacacta 300
     298 cagttgacaa aagttaatct cggtgataag aatatgc
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     306 <223> OTHER INFORMATION: Genbank Accession No. AA026641
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Use of n and/or Xaa has been detected in the Sequence Listing. Review the Sequence Listing to insure a corresponding explanation is presented in the <220> to <223> fields of each sequence using n or Xaa.

VERIFICATION SUMMARY

6

DATE: 12/07/2001 TIME: 19:16:26

PATENT APPLICATION: US/09/960,706

Input Set : N:\jumbos\960706.txt

Output Set: N:\CRF3\12072001\I960706.raw

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L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:58 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2
L:59 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2
L:114 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5
L:116 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5
L:139 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6
L:169 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:171 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:202 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10
L:206 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10
L:243 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12
L:244 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12
L:281 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14
L:314 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:16
L:315 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:16
L:318 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:16
L:319 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:16
L:320 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:16
L:358 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:18
L:359 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:18
L:360 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:18
L:394 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:20
L:395 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:20
L:414 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21
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L:564 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28
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L:663 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:33
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VERIFICATION SUMMARY

DATE: 12/07/2001

PATENT APPLICATION: US/09/960,706

TIME: 19:16:26

Input Set : N:\jumbos\960706.txt

Output Set: N:\CRF3\12072001\1960706.raw

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DATE: 12/06/2001 RAW SEQUENCE LISTING TIME: 09:44:18 PATENT APPLICATION: US/09/960,706

Input Set : N:\jumbos\960706.txt

Output Set: N:\CRF3\12062001\I960706.raw

gus

Does Not Comply

Corrected Diskette Needed

3 <110> APPLICANT: Munger, William E. 5 <120> TITLE OF INVENTION: Identifying Drugs for and Diagnosis of Benign Prostatic Hyperplasia Using

Gene Expression Profiles

8 <130> FILE REFERENCE: 44921-5029-01US

C--> 10 <140> CURRENT APPLICATION NUMBER: US/09/960,706

C--> 11 <141> CURRENT FILING DATE: 2001-09-24

13 <150> PRIOR APPLICATION NUMBER: 60/223,323

14 <151> PRIOR FILING DATE: 2000-08-07

16 <150> PRIOR APPLICATION NUMBER: 09/873,319

17 <151> PRIOR FILING DATE: 2001-06-05

19 <160> NUMBER OF SEQ ID NOS: 1124

21 <170> SOFTWARE: PatentIn Ver. 2.1

ERRORED SEQUENCES

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33793 gttttgtttt tttgggggag tattatttct attatttatg ttttagttat aggacgcgtg 540

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VERIFICATION SUMMARY
PATENT APPLICATION: US/09/960,706

DATE: 12/06/2001 TIME: 09:44:25

Input Set : N:\jumbos\960706.txt

Output Set: N:\CRF3\12062001\I960706.raw

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L:690 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:34



VERIFICATION SUMMARY

PATENT APPLICATION: US/09/960,706

DATE: 12/06/2001 TIME: 09:44:25

Input Set : N:\jumbos\960706.txt

Output Set: N:\CRF3\12062001\I960706.raw

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